

From: [Robert Neely](#)
To: [Eric Blischke/R10/USEPA/US@EPA](#)
Subject: Fw: Willamette River watershed database posted - 07/30/08
Date: 08/13/2008 06:26 PM

Fyi

----- Original Message -----

From: Dan Rutz <danr@genwest.com>
To: Robert.Neely@noaa.gov <Robert.Neely@noaa.gov>; Jay Field <Jay.Field@noaa.gov>
Cc: cgsevern@premiercorp-usa.com <cgsevern@premiercorp-usa.com>; 'Benjamin Shorr' <Benjamin.Shorr@noaa.gov>; blischke.eric@epa.gov <blischke.eric@epa.gov>
Sent: Fri Aug 01 17:06:38 2008
Subject: Re: Willamette River watershed database posted - 07/30/08

Hi everyone,

Please find the July 29 Willamette River data and corresponding dictionaries posted to response.restoration.noaa.gov/querymanager.

Dan

P.S. The IC reports for the three datasets can be found at:

Non-Responsive

P.P.S. The IC flagged one new item (the same item in all three data sets):

Chem
Fields tested, What was tested, Test results (1)
ChemCode+Units, RELATION INTO ChemDict, SPEC_GRAV NA

From: Robert.Neely@noaa.gov
To: Jay Field [mailto:Jay.Field@noaa.gov]
Cc: cgsevern@premiercorp-usa.com, 'Dan Rutz' [mailto:danr@genwest.com], 'Benjamin Shorr' [mailto:Benjamin.Shorr@noaa.gov], blischke.eric@epa.gov
Sent: Wed, 30 Jul 2008 10:19:36 -0700
Subject: Re: Willamette River watershed database posted - 07/30/08

Let's see what Eric learns once he's discussed with LWG.

-R

----- Original Message -----

From: Jay Field <Jay.Field@noaa.gov>
Date: Wednesday, July 30, 2008 10:12 am
Subject: Re: Willamette River watershed database posted - 07/30/08

> Rob,
> if we're expecting resolution of the carp/bass data in the near
> term
> (next couple of weeks), I recommend waiting until those issues are
> resolved.
> Jay
>
> Robert.Neely@noaa.gov wrote:
> > Hey -- Thanks Corinne. I believe Jay is expecting the stats for
> the bioassays from LWG soon.
> > On the R3 tissue data, turns out there are some potential issues
> with the carp and bass data
> > that EPA is working to resolve with LWG. (Analyses were done on
> whole body and filets for
> > these receptors and it's not clear that the filet results were
> mathematically recombined with
> > the whole body.) So for now we should hold on doing the bass and
> carp, but the others
> > (crayfish, sculpin, etc.) should be OK. Does this create any
> efficiency problems or
> > complications that would make it better to weight until all R3
> tissue issues are resolved, or
> > can we move ahead with the ones that are ready?

> > -R

> > ----- Original Message -----

> > From: Corinne Severn <cgsevern@premiercorp-usa.com>
> > Date: Wednesday, July 30, 2008 9:51 am
> > Subject: Willamette River watershed database posted - 07/30/08

> > Jay/Rob/Ben/Dan -

> > I have posted three new Willamette River files to the NOAA FTP
> > server in the
> > folder named as follows:
> > /ARD/Query_Manager/Data_Files_zipped/Willamette/

> > The files have the following names:

> > WR080729.zip - This file contains all data
> > WRCat1_080729.zip - This file (Nature and Extent) contains any
> > data
> > with "Cat1" in the chemcat field
> > WRQA2Cat1_080729.zip - This file (Risk) contains only data with

> >> QA2Cat1classification.
> >>
> >> There is also a new dictionary file (Dict080729.zip) which
> contains
> >> newestIDs required for the bioassay studies.
> >>
> >> To the latest version, we have added the Round 3 data from the
> >> sedimentcores, co-located sediments (which will correspond to
> >> tissue data to be
> >> added shortly), and the Round 3 bioassay data with associated
> sediment>> chemistry.
> >>
> >> For the bioassay data, there are several items to be aware of -
> >>
> >> 1) Currently, there are no results available for the statistic
> >> significancefor the data. For the interim, we have marked those
> >> results where the
> >> control-adjusted value is <80% as significant.
> >>
> >> 2) The tests results for growth for Chironomus dilutus are based
> on
> >> theaverage individual ash-free biomass in mg from the original
> >> laboratoryreports (ie., total ash-free dry weight/count of
> animals
> >> at end of test
> >> coded as AFDW in lab files).
> >> The test results for growth for Hyalella azteca are based on the
> >> averageindividual biomass in mg from the original laboratory
> >> reports (ie., total
> >> dry weight/count of animals at end of test coded as WT in lab
> >> files).>> These are the same endpoints used for the Round 2
> bioassay data for
> >> thegrowth endpoints.
> >>
> >> 3) An additional growth endpoint was coded from the laboratory
> >> bioassay data
> >> based on total biomass in mg (TWT for H. azteca and TAFDW for C.
> >> dilutus in
> >> the original lab files). This result is not divided by the
> >> animals, but is
> >> the sum of all organism weights for replicate bioassay results
> >> (biorep.dbftable) or the average of the total organism weights
> for
> >> all replicates for
> >> the mean bioassay results (biosumm.dbf table).
> >>
> >> Thanks,
> >>
> >> Corinne
> >>
> >>
> >> _____
> >> Corinne Severn
> >> Premier Environmental Services
> >> Ph: 702-255-9685
> >> Fax: 888-220-9867
> >> Cell: 206-226-9663
> >>
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